

SEQUENCE LISTING

<110> BAYER AG

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<170> PatentIn Ver. 2.1

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<222> (244)..(1566)

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120

gatagactct agtggatatc tacaagtatt gtttttttga taaaataggc tgaggtgaga
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aggttaacata aaggaaagac aaaaacttgg gaattgtttt agaccaccga ggtttcttgt
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288

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336

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cgc gtc ttt gta tca tcc agg ttc cta gct cga gat agg aat ttg atg
384

Arg Val Phe Val Ser Ser Arg Phe Leu Ala Arg Asp Arg Asn Leu Met
35 40 45

tgg aat ggg aga atc aag aaa ggt ggg aga caa agg tgg aat ttt ggc
432

Trp Asn Gly Arg Ile Lys Lys Gly Gly Arg Gln Arg Trp Asn Phe Gly
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480

Ser Leu Ile Ala Asp Pro Arg Tyr Ser Cys Leu Gly Gly Ser Arg Thr
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gaa aag gga agc act ttc tct gta cag tcc agt ttg gtg gct agc cca
 528

Glu Lys Gly Ser Thr Phe Ser Val Gln Ser Ser Leu Val Ala Ser Pro
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gct gga gaa atg act gtg tca tca gag aaa aag gtg tat gat gtg gta
 576

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tta aag cag gca gct tta gtg aag agg cag ctg aga tct acc gat gat
 624

Leu Lys Gln Ala Ala Leu Val Lys Arg Gln Leu Arg Ser Thr Asp Asp
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 720

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Ile Trp Ala Ile Tyr Val Trp Cys Arg Arg Thr Asp Glu Leu Val Asp
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 912

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Ala Ala Leu Ser Asp Thr Val Ser Arg Phe Pro Val Asp Ile Gln Pro

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Lys Gln Ala Ala Leu Val Lys Arg Gln Leu Arg Ser Thr Asp Asp Leu
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240
aaattaagca gaggaaggaa ggaaacagaa aacagaaagt aagacaaaaa accttggaaat
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Gly Ser Val Arg Ser Ala Met Val Ala Thr Pro Ala Gly Glu Met Ala
60 65 70
acg atg aca tca gaa cag atg gtt tat gat gtg gtt tta aaa caa gca
593
Thr Met Thr Ser Glu Gln Met Val Tyr Asp Val Val Leu Lys Gln Ala
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Ala Leu Val Lys Arg Gln Leu Arg Ser Ala Asp Asp Leu Glu Val Lys
90 95 100

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 737

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Gly Thr Met Xaa Met Thr Pro Glu Arg Arg Arg Ala Ile Trp Ala Ile
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Tyr Val Trp Cys Arg Arg Thr Asp Glu Leu Val Asp Gly Pro Asn Ala
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Ser His Ile Thr Pro Gln Ala Leu Asp Arg Trp Glu Asp Arg Leu Glu
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 929

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 977

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 1169

DRAFT
REVIEWED
RECORDED
TO FILE

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1313

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345 350 355

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360 365 370 375

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Phe Thr Lys Arg Ala Tyr Val Ser Lys Pro Lys Lys Leu Ile Ser Leu
380 385 390

cct att gct tat gca aaa tct ctt gtg ccc cct aca aga act ctt gtc
1553

Pro Ile Ala Tyr Ala Lys Ser Leu Val Pro Pro Thr Arg Thr Leu Val
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1605

Thr Ser Ser
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Ala Asp Asp Leu Glu Val Lys Pro Glu Ile Pro Leu Pro Gly Asn Leu
100 105 110

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115 120 125

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Arg Arg Ala Ile Trp Ala Ile Tyr Val Trp Cys Arg Arg Thr Asp Glu
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165 170 175

Arg Trp Glu Asp Arg Leu Glu Asp Val Phe Ser Gly Arg Pro Phe Asp
180 185 190

Met Leu Asp Ala Ala Leu Ser Asp Thr Val Ser Lys Phe Pro Val Asp
195 200 205

Ile Gln Pro Phe Arg Asp Met Ile Glu Gly Met Arg Met Asp Leu Arg
210 215 220

Lys Ser Arg Tyr Arg Asn Phe Asp Glu Leu Tyr Leu Tyr Cys Tyr Tyr
225 230 235 240

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245 250 255

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260 265 270

Leu Gly Ile Ala Asn Gln Leu Thr Asn Ile Leu Arg Asp Val Gly Glu
275 280 285

Asp Ala Arg Arg Gly Arg Val Tyr Leu Pro Gln Asp Glu Leu Ala Gln
290 295 300

Ala Gly Leu Phe Asp Asp Asp Ile Phe Ala Gly Lys Val Thr Asp Lys
305 310 315 320

Trp Arg Ser Phe Met Lys Lys Gln Ile Gln Arg Ala Arg Lys Phe Phe
325 330 335

Asp Glu Ala Glu Glu Gly Val Thr Gln Leu Ser Ser Ala Ser Arg Trp
340 345 350

Pro Val Trp Ala Ser Leu Leu Leu Tyr Arg Gln Ile Leu Asp Glu Ile
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Glu Ala Asn Asp Tyr Asn Asn Phe Thr Lys Arg Ala Tyr Val Ser Lys
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120

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180

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230

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422		
Ala Pro Lys Gly Leu Phe Pro Pro Glu Pro Glu His Tyr Arg Gly Pro		
65	70	75
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470		
Lys Leu Lys Val Ala Ile Ile Gly Ala Gly Leu Ala Gly Met Ser Thr		
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518		
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Arg Pro Phe Ile Gly Gly Lys Val Gly Ser Phe Val Asp Arg Arg Gly		
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145	150	155
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195	200	205

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998

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1046

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1094

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1142

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305 310 315

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1190

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320 325 330

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1238

Met Tyr Val Ser Gly Leu Ala Met Ser Lys Ala Thr Gln Lys Lys Val
335 340 345 350

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1286

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355 360 365

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1334

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 Ser Val Val Lys Ile Gly Gln Ser Leu Tyr Arg Glu Gly Pro Gly Lys
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1910

Lys Leu Val Val Leu Arg Lys Lys Ile Ala Ala Ala Glu Ser Asn Glu
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1955

Ile Ser Glu Gly Val Ser Val Ser Asp Glu Leu Ser Leu Val
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Lys Gly Leu Phe Pro Pro Glu Pro Glu His Tyr Arg Gly Pro Lys Leu
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Lys Val Ala Ile Ile Gly Ala Gly Leu Ala Gly Met Ser Thr Ala Val
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Glu Leu Leu Asp Gln Gly His Glu Val Asp Ile Tyr Glu Ser Arg Pro
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Phe Ile Gly Gly Lys Val Gly Ser Phe Val Asp Arg Arg Gly Asn His
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 Ile Glu Met Gly Leu His Val Phe Phe Gly Cys Tyr Asn Asn Leu Phe
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 His Thr His Thr Phe Val Asn Lys Gly Gly Glu Ile Gly Glu Leu Asp
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 Phe Arg Phe Pro Val Gly Ala Pro Leu His Gly Ile Asn Ala Phe Leu
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 Ser Thr Asn Gln Leu Lys Ile Tyr Asp Lys Ala Arg Asn Ala Val Ala
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 Met Ser Lys Gly Gly Thr Arg Ala Ser Ile Gln Arg Met Trp Asp Pro
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 Gly Cys Arg Gln Val Leu Tyr Glu Thr Ser Ser Asp Gly Ser Met Tyr
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 Leu Val Gly Val Pro Val Val Thr Val Gln Leu Arg Tyr Asn Gly Trp
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 Thr Gly Leu Asp Asn Leu Leu Tyr Thr Pro Asp Ala Asp Phe Ser Cys
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Gln Gly Ser Leu Leu Gln Cys Val Leu Thr Pro Gly Asp Pro Tyr Met		
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Pro Leu Leu Asn Asp Glu Ile Ile Lys Arg Val Ser Lys Gln Val Leu		
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Ala Leu Phe Pro Ser Ser Gln Gly Leu Glu Val Thr Trp Ser Ser Val		
485	490	495
Val Lys Ile Gly Gln Ser Leu Tyr Arg Glu Gly Pro Gly Lys Asp Pro		
500	505	510
Phe Arg Pro Asp Gln Lys Thr Pro Val Glu Asn Phe Phe Leu Ala Gly		
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Ser Tyr Thr Lys Gln Asp Tyr Ile Asp Ser Met Glu Gly Ala Thr Leu		
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Ser Gly Arg Gln Ala Ser Ala Tyr Val Cys Asp Ala Gly Glu Lys Leu		
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23

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